Docket No.: CHIR-0337

App No.: Not Yet Assigned Filed: Herewith Title: HELICOBACTER PYLORI CAI ANTIGEN Inventors: Covacci et al.

Attorney: Felicity E. Groth Phone: (215) 568

Sheet 1 of 14

Phone: (215) 568-3100

1/14

					1/14
1	AAAAAGAAAG	GAAGAAAATG	GAAATACAAC	AAACACACCG	CAAAATCAAT
51	CGCCCTCTGG	TTTCTCTCGC	TTTAGTAGGA	GCATTAGTCA	GCATCACACC
101	GCAACAAAGT	CATGCCGCCT	TTTTCACAAC	CGTGATCATT	CCAGCCATTG
1,51	TTGGGGGTAT	CGCTACAGGC	ACCGCTGTAG	GAACGGTCTC	AGGGCTTCTT
201	AGCTGGGGGC	TCAAACAAGC	CGAAGAAGCC	AATAAAACCC	CAGATAAACC
251	CGATAAAGTT	TGGCGCATTC	AAGCAGGAAA	AGGCTTTAAT	GAATTCCCTA
301	ACAAGGAATA	CGACTTATAC	AGATCCCTTT	TATCCAGTAA	GATTGATGGA
351	GGTTGGGATT	GGGGGAATGC	CGCTAGGCAT	TATTGGGTCA	AAGGCGGGCA
401	ACAGAATAAG	CTTGAAGTGG	ATATGAAAGA	CGCTGTAGGG	ACTTATACCT
451	TATCAGGGCT	TAGAAACTTT	ACTGGTGGGG	ATTTAGATGT	CAATATGCAA
501	AAAGCCACTT	TACGCTTGGG	CCAATTCAAT	GGCAATTCTT	TTACAAGCTA
551	TAAGGATAGT	GCTGATCGCA	CCACGAGAGT	GATTTCAACG	CTAAAAATAT
601	CTCAATTGAT	AATTTTGCAG	AAATCAACAA	CTCGTGTGGG	TTCTGGAGCC
651	GGGAGGAAAG	CCAGCTCTAC	GGTTTTGACT	TTGCAAGCTT	CAGAAGGGAT
701	CACTAGCGAT	AAAAACGCTG	AAATTTCTCT	TTATGATGGT	GCCACGCTCA
751	ATTTGGCTTC	AAGCAGCGTT	AAATTAATGG	GTAATGTGTG	GATGGGCCGT
801	TTGCAATACG	TGGGAGCGTA	TTTGGCCCCT	TCATACAGCA	CGATAAACAC
851	TTCAAAAGTA	ACAGGGGAAG	TGAATTTTAA	CCACCTCACT	GTTGGCGATA
901	AAAACGCCGC	TCAAGCGGGC	ATTATCGCTA	ATAAAAAGAC	TAATATTGGC
951	ACACTGGATT	TGTGGCAAAG	CGCCGGGTTA	AACATTATCG	CTCCTCCAGA
1001	AGGTGGCTAT	AAGGATAAAC	CCAATAATAC	CCCTTCTCAA	AGTGGTGCTA
1051	AAAACGACAA	AAATGAAAGC	GCTAAAAACG	ACAAACAAGA	GAGCAGTCAA
1101	AATAATAGTA	ACACTCAGGT	CATTAACCCA	CCCAATAGTG	CGCAAAAAAC
1151	AGAAGTTCAA	CCCACGCAAG	TCATTGATGG	GCCTTTTGCG	GGCGGCAAAG
1201	ACACGGTTGT	CAATATCAAC	CGCATCAACA	CTAACGCTGA	TGGCACGATT
1251	AGAGTGGGAG	GGTTTAAAGC	TTCTCTTACC	ACCAATGCGG	CTCATTTGCA
1301	TATCGGCAAA	GGCGGTGTCA	ATCTGTCCAA	TCAAGCGAGC	GGGCGCTCTC
	•	1			

FIG. 1A

Docket No.: CHIR-0337
App No.: Not Yet Assigned Filed: Herewith
Title: HELICOBACTER PYLORI CAI ANTIGEN
Inventors: Covacci et al.
Attomey: Felicity E. Groth Phone: (215) 568
Sheet 2 of 14

Phone: (215) 568-3100

					2/14
1351	TTATAGTGGA	AAATCTAACT	GGGAATATCA	CCGTTGATGG	GCCTTTAAGA
1401	GTGAATAATC	AAGTGGGTGG	CTATGCTTTG	GCAGGATCAA	GCGCGAATTT
1451	TGAGTTTAAG	GCTGGTACGG	ATACCAAAAA	CGGCACAGCC	ACTTTTAATA
1501	ACGATATTAG	TCTGGGAAGA	TTTGTGAATT	TAAAGGTGGA	TGCTCATACA
1551	GCTAATTTTA	AAGGTATTGA	TACGGGTAAT	GGTGGTTTCA	ACACCTTAGA
1601	TTTTAGTGGC	GTTACAGACA	AAGTCAATAT	CAACAAGCTC	ATTACGGCTT
1651	CCACTAATGT	GGCCGTTAAA	AACTTCAACA	TTAATGAATT	GATTGTTAAA
1701	ACCAATGGGA	TAAGTGTGGG	GGAATATACT	CATTTTAGCG	AAGATATAGG
1751	CAGTCAATCG	CGCATCAATA	CCGTGCGTTT	GGAAACTGGC	ACTAGGTCAC
1801	TTTTCTCTGG	GGGTGTTAAA	TTTAAAGGTG	GCGAAAAATT	GGTTATAGAT
1851	GAGTTTTACT	ATAGCCCTTG	GAATTATTTT	GACGCTAGAA	ATATTAAAAA
1901	TGTTGAAATC	ACCAATAAAC	TTGCTTTTGG	ACCTCAAGGA	AGTCCTTGGG
1951	GCACATCAAA	ACTTATGTTC	AATAATCTAA	CCCTAGGTCA	AAATGCGGTC
2001	ATGGATTATA	GCCAATTTTT	AAATTTAACC	ATTCAAGGGG	ATTTCATCAA
2051	CAATCAAGGC	ACTATCAACT	ATCTGGTCCG	AGGTGGGAAA	GTGGCAACCT
2101	TAAGCGTAGG	CAATGCAGCA	GCTATGATGT	TTAATAATGA	TATAGACAGC
2151	GCGACCGGAT	TTTACAAACC	GCTCATCAAG	ATTAACAGCG	CTCAAGATCT
2201	CATTAAAAAT	ACAGAACATG	TTTTATTGAA	AGCGAAAATC	ATTGGTTATG
2251	GTAATGTTTC	TACAGGTACC	AATGGCATTA	GTAATGTTAA	TCTAGAAGAG
2301	CAATTCAAAG	AGCGCCTAGC	CCTTTATAAC	AACAATAACC	GCATGGATAC
2351	TTGTGTGGTG	CGAAATACTG	ATGACATTAA	AGCATGCGGT	ATGGCTATCG
2401	GCGATCAAAG	CATGGTGAAC	AACCCTGACA	ATTACAAGTA	TCTTATCGGT
2451	AAGGCATGGA	AAAATATAGG	GATCAGCAAA	ACAGCTAATG	GCTCTAAAAT
2501	TTCGGTGTAT	TATTTAGGCA	ATTCTACGCC	TACTGAGAAT	GGTGGCAATA
2551	CCACAAATTT	ACCCACAAAC	AGCACTAGCA	ATGCACGTTC	TGCCAACAAC
2601	GCCCTTGCAC	AAAACGCTCC	TTTCGCTCAA	CCTAGTGCTA	CTCCTAATTT
2651	AGTCGCTATC	AATCAGCATG	ATTTTGGCAC	TATTGAAAGC	GTGTTTGAAT

3/14

					3/14
2701	TGGCTAACCG	CTCTAAAGAT	ATTGACACGC	TTTATGCTAA	CTCAGGCGCT
2751	CAAGGCAGGG	ATCTCTTACA	AACCTTATTG	ATTGATAGCC	ATGATGCGGG
2801	TTATGCCAGA	AAAATGATTG	ATGCTACAAG	CGCTAATGAA	ATCACCAAGC
2851	AATTGAATAC	GGCCACTACC	ACTTTAAACA	ACATAGCCAG	TTTAGAGCAT
2901	AAAACCAGCG	GCTTACAAAC	TTTGAGCTTG	AGTAATGCGA	TGATTTTAAA
2951	TTCTCGTTTA	GTCAATCTCT	CCAGGAGACA	CACCAACCAT	ATTGACTCGT
3001	TCGCCAAACG	CTTACAAGCT	TTAAAAGACC	AAAAATTCGC	TTCTTTAGAA
3051	AGCGCGGCAG	AAGTGTTGTA	TCAATTTGCC	CCTAAATATG	AAAAACCTAC
3101	CAATGTTTGG	GCTAACGCTA	TTGGGGGAAC	GAGCTTGAAT	AATGGCTCTA
3151	ACGCTTCATT	GTATGGCACA	AGCGCGGGCG	TAGACGCTTA	CCTTAACGGG
3201	CAAGTGGAAG	CCATTGTGGG	CGGTTTTGGA	AGCTATGGTT	ATAGCTCTTT
3251	TAATAATCGT	GCGAACTCCC	TTAACTCTGG	GGCCAATAAC	ACTAATTTTG
3301	GCGTGTATAG	CCGTATTTTA	ACCAACCAGC	ATGAATTTGA	CTTTGAAGCT
3351	CAAGGGCAC	TAGGGAGCGA	TCAATCAAGC	TTGAATTTCA	AAAGCGCTCT
3401	ATTACAAGAT	TTGAATCAAA	GCTATCATTA	CTTAGCCTAT	AGCGCTGCAA
3451	CAAGAGCGAG	CTATGGTTAT	GACTTCGCGT	TTTTTAGGAA	CGCTTTAGTG
3501	TTAAAACCAA	GCGTGGGTGT	GAGCTATAAC	CATTTAGGTT	CAACCAACTT
3551	TAAAAGCAAC	AGCACCAATC	AAGTGGCTTT	GAAAAATGGC	TCTAGCAGTC
3601	AGCATTTATT	CAACGCTAGC	GCTAATGTGG	AAGCGCGCTA	TTATTATGGG
3651	GACACTTCAT	ACTTCTACAT	GAATGCTGGA	GTTTTACAAG	AGTTCGCTCA
3701	TGTTGGCTCT	AATAACGCCG	CGTCTTTAAA	CACCTTTAAA	GTGAATGCCG
3751	CTCGCAACCC	TTTAAATACC	CATGCCAGAG	TGATGATGGG	TGGGGAATTA
3801	AAATTAGCTA	AAGAAGTGTT	TTTGAATTTG	GGCGTTGTTT	ATTTGCACAA
3851	TTTGATTTCC	AATATAGGCC	ATTTCGCTTC	CAATTTAGGA	ATGAGGTATA
3901	GTTTCTAAAT	ACCGCTCTTA	AACCCATGCT	CAAAGCATGG	GTTTGAAATC
3951	TTACAAAACA	 ,	0 40		
		 1	G 1C		

FIG. 1C

Docket No.: CHIR-0337
App No.: Not Yet Assigned Filed: Herewith

Title: HELICOBACTER PYLORI CAI ANTIGEN

Inventors: Covacci et al. Attorney: Felicity E. Groth Sheet 4 of 14

Phone: (215) 568-3100

4/14 1 MEIQQTHRKI NRPLVSLALV GALVSITPQQ SHAAFFTTVI IPAIVGGIAT 51 GTAVGTVSGL LSWGLKQAEE ANKTPDKPDK VWRIQAGKGF NEFPNKEYDL 101 YRSLLSSKID GGWDWGNAAR HYWVKGGQQN KLEVDMKDAV GTYTLSGLRN 151 FTGGDLDVNM QKATLRLGQF NGNSFTSYKD SADRTTRVIS TLKISQLIIL 201 QKSTTRVGSG AGRKASSTVL TLQASEGITS DKNAEISLYD GATLNLASSS 251 VKLMGNVWMG RLQYVGAYLA PSYSTINTSK VTGEVNFNHL TVGDKNAAQA 301 GIIANKKTNI GTLDLWQSAG LNIIAPPEGG YKDKPNNTPS QSGAKNDKNE 351 SAKNDKQESS QNNSNTQVIN PPNSAQKTEV QPTQVIDGPF AGGKDTVVNI 401 NRINTNADGT IRVGGFKASL TTNAAHLHIG KGGVNLSNQA SGRSLIVENL 451 TGNITVDGPL RVNNQVGGYA LAGSSANFEF KAGTDTKNGT ATFNNDISLG 501 RFVNLKVDAH TANFKGIDTG NGGFNTLDFS GVTDKVNINK LITASTNVAV 551 KNFNINELIV KTNGISVGEY THFSEDIGSQ SRINTVRLET GTRSLFSGGV 601 KFKGGEKLVI DEFYYSPWNY FDARNIKNVE ITNKLAFGPQ GSPWGTSKLM 651 FNNLTLGQNA VMDYSQFLNL TIQGDFINNQ GTINYLVRGG KVATLSVGNA 701 AAMMFNNDID SATGFYKPLI KINSAQDLIK NTEHVLLKAK IIGYGNVSTG 751 TNGISNVNLE EQFKERLALY NNNNRMDTCV VRNTDDIKAC GMAIGDQSMV 801 NNPDNYKYLI GKAWKNIGIS KTANGSKISV YYLGNSTPTE NGGNTTNLPT 851 NTTSNARSAN NALAQNAPFA QPSATPNLVA INQHDFGTIE SVFELANRSK 901 DIDTLYANSG AGGRDLLQTL LIDSHDAGYA RKMIDATSAN EITKQLNTAT 951 TTLNNIASLE HKTSGLQTLS LSNAMILNSR LVNLSRRHTN HIDSFAKRLQ 1001 ALKDOKFASL ESAAEVLYQF APKYEKPTNV WANAIGGTSL NNGSNASLYG 1051 TSAGVDAYLN GQVEAIVGGF GSYGYSSFNN RANSLNSGAN NTNFGVYSRI 1101 LTNQHEFDFE AQGALGSDQS SLNFKSALLQ DLNQSYHYLA YSAATRASYG 1151 YDFAFFRNAL VLKPSVGVSY NHLGSTNFKS NSTNQVALKN GSSSQHLFNA

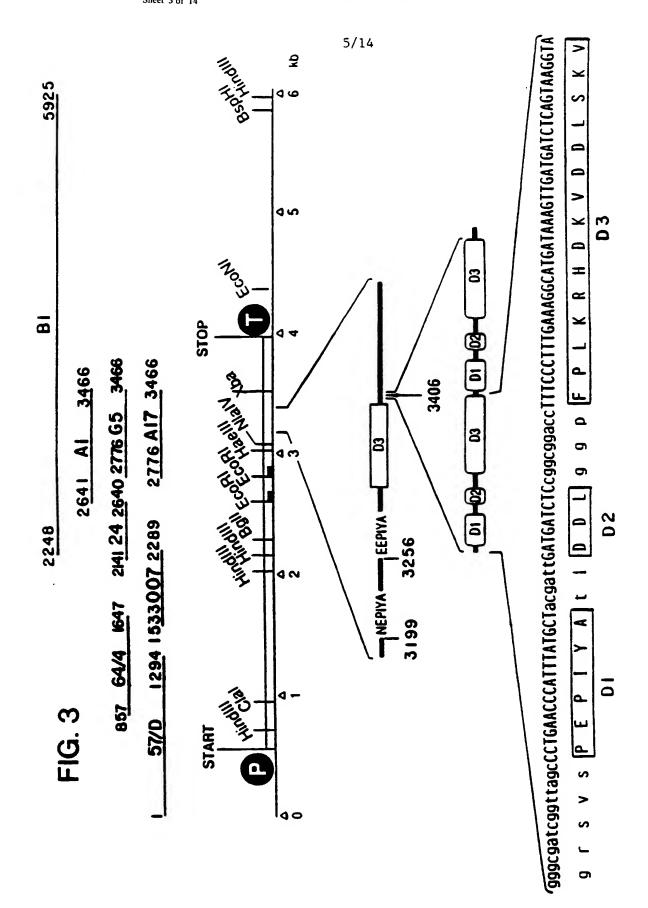
1251 THARVMMGGE LKLAKEVFLN LGVVYLHNLI SNIGHFASNL GMRYSF

1201 SANVEARYYY GDTSYFYMNA GVLQEFAHVG SNNAASLNTF KVNAARNPLN

Docket No.: CHIR-0337 App No.: Not Yet Assigned Filed: Herewith Title: HELICOBACTER PYLORI CAI ANTIGEN Inventors: Covacci et al. Attomey: Felicity E. Groth Phone: (215) 56

Sheet 5 of 14

Phone: (215) 568-3100



Inventors: Covacci et al. Attorney: Felicity E. Groth Sheet 6 of 14

Phone: (215) 568-3100

6/14

CTCCATTTTAAGCAACTCCATAGACCACTAAAGAAACTTTTTTTGAGGCTATCTTTGAAA GCTTAATTATACATGCTATAGTAAGCATGACACACACACCAAACTATTTTTAGAACGCTT TCAAAAAGATTCATTTCTTATTTCTTGTTCTTATTAAAGTTCTTTCATTTTAGCAAATTT TTGACACTAACAAGATACCGATAGGTATGAAACTAGGTATAGTAAGGAGAAACAATGACT AATAATCTTCAAGTAGCTTTTCTTAAAGTTGATAACGCTGTCGCTTCATACGATCCTGAT 23 N N L Q V A F L K V D N A V A SYD CAATTAAGGGAAGAATACTCCAATAAAGCGATCAAAAATCCTACCAAAAAGAATCAGTAT 63 Q L R E E Y S N K A I K N P T K K N Q Y GAATCTTCCACAAAGAGCTTTCAGAAATTTGGGGATCAGCGTTACCGAATTTTCACAAGT 103 E S S T K S F Q K F G D Q R Y R I F T S GAAAATATCATACAACCCCCTATCCTTGATGATAAAGAGAAAGCGGAGTTTTTGAAATCT NIIQPPILDDKEKAEFL ATGGGCGTGTTTGATGAGTCCTTGAAAGAAAGGCAAGAAGCAGAAAAAAATGGAGAGCCT 183 M G V F D E S L K E R Q E A E K N G E P GATGTCAAAGAAGCAATCAATCAAGAACCAGTTCCCCATGTCCAACCAGATATAGCCACT 223 D V K E A I NQEPVPHVQPDI AATTTTTCTAAATTCACTCTTGGCGATATGGAAATGTTAGATGTTGAGGGAGTCGCTGAC 263 N F S K F T L G D M E M L D V E G V A TTAATGGGGAGTCATAATGGCATAGAACCTGAAAAAGTTTCATTGTTGTATGGGGGCAAT 303 L M G S H N G I E P E K V SLLYGGN AACAATGTGGCTACAATAATTAATGTGCATATGAAAAACGGCAGTGGCTTAGTCATAGCA 343 N N V A T I I NVHMKNGS G L GGCTCACAACGAGCATTAAGTCAAGAAGAGATCCAAAACAAAATAGATTTCATGGAATTT 383 G S Q- R A L S Q E E I Q N K I DFMEF **ACTGAGATTAAAGATTTCCAAAAAGACTCTAAGGCTTATTTAGACGCCCTAGGGAATGAT** 423 TEIKDFQKDSKAYLD A L AATGGGGATTTGAGCTACACTCTCAAAGATTATGGGAAAAAAGCAGATAAAGCTTTAGAT 463 N G D L S Y T L K D Y G K K A D K A L D

FIG. 4A

TATTCTAATTCAAATACACCAACGCCTCCAAGAATCCCAATAAGGGTGTAGGCGTTACG

A	TCTO	STCC	TAT	TGA	TTT	GTT	TTO	CAT	TTT	GTT	TTC(CCAT	GT	3GA1	CT.	TGT	GGAT	CAC	CAAAC	120
																			TTTCT	240
T.	TGTT	AAT	TGT	GGG	TAA	AAA	TGT	TGAA	TCG	TCC	CTAG	SCC1	TT	AGAC	GC	CTG	CAAC	GAT	CGGG	360
A	ATGA	GAA	TGT	TCA	AAG	ACA	TGA	ATT	GAC	TAC	TCA	AAGO	GTO	STAG	GCG/	ATT	TTTA	\GC#	AGTCT	480
																			TATC	600
N	Ε	T	I	D	Q	Q	P	Q	T	Ε	A	Α	F	N	Ρ	Q	Q	F	1	
C	AAAA	ACC	AAT	CGT	TGA	TAA	GAA	CGA	TAG	GGA	TAA	CAG	GCA	AGC	TT	TTG/	AAGG	AAT	CTCG	720
Q	Κ	Ρ	I	٧	D	K	N	D	R	D	N	R	Q	Α	F	Ε	G	Ī	S	
T	TTTC	AGA	CTT	TAT	CAA	TAA	GAG	CAA	TĠA	TTT	'AAT	CAA	CAA	AGA	CA	ATCI	CAT	TGA	TGTA	840
F	S	D	F	I	N	K	S	N	D	L	I	N	K	D	N	L	I	D	V	.
TO	GGT	GTC	CCA	TCA	AAA	CGA	TCC	GTC	TAA	AAT	CAA	CAC	CCG	ATC	GAT	CCC	SAAA	TTT	TATG	960
W	٧	S	Н	Q	N	D	Ρ	S	K	I	N	T	R	S	I	R	N	F	M	
GC	CAA	ACA.	ATC'	TTT	TGC	AGG	AAT	CAT	TAT	AGG	GAA	TCA	AAT	CCG	AAC	:GG/	ATCA	AAA	GTTC	1080
Α	K	Q	S	F	A	G	I	I	I	G	N	Q	I	R	T	D	Q	K	F	
AC	TGG	TGG	GGA'	TTG	GTT	GGA	TAT	TTT	TCT	CTC	ATT	TAT	ATT	TGA	CAA	AAA	ACA	ATC	TTCT	1200
Т	G	G	D	W	L	D	I	F	L		F	Ī	F	D	K	K	Q	S	S	1200
AC	CAC	CAC	CGA	CAT	ACA	AGG	CTT	ACC	GCC			TAG	AGA	TTT			_	-	GGGT	1320
Т	T	T	D	Ī	Q	G	L	Р	P	E	A	R	D	L	L	D	E	R	G .	1720
ΑT	TGA	TCC	CAA'	TTA	CAA	•	CAA	-		_			_	TAA	CGC	_	_		TGTG	1440
Ī	D	ρ	N	Υ	K	F	N	Q	L	L	I	Н	N	N	A	1	S	S	v	1440
GG	TGG	TCC:	TGG	AGC1		GCA		_	_	CGC	-					AGA	_	_	AGGC	1560
G	G	Р	G	A	R	Н	D	W	N	A	T	V	G	Υ	K	D	Q	Q	G	1500
GG	_	-	-		• •		_				•	-	_	-		_	_	_	CACA	1680
G	G	E	K	G	1	N	N	Р	S	F	Υ	L	Υ	K	Ε	D	Q	L	T	1000
_	_	_	• •	-	•			-	-	-	-	_	-		_	_	_	-	CCGA	1800
1	Α.	Q		N	A	K	1	D	N	L	S	E	K	E	K	E	K	F	R	1000
CG		_	• •		••		_	_		_	_	_		_		_		•	TGGT	1920
R	1	A	F	V	S	K	K	D	T	K	H	S	A	1	7	T	E	F	G	1320
. •	•	• •	•	•	•			_	•			•	•••	L TGG	Cgt	'GAT	_	•	TGAT	2040
R	E	K	N	V .	T	1	Q	AGG G	S			H	D	G	V	M	F	V	D	2040
•	_		••	•	•	•	_	-	-	_		• •	•	•	-	• •	•	•	TGAT	2150
A A	וטטי	CUI	116	LHI	111	NOP	וטא	יטטא	CII	I AA	CAA	וטטו	HUL	IAI	CII	IAA	III.	ULL	IGAI	2160

FIG. 4B

Inventors: Covacci et al. Attorney: Felicity E. Groth Sheet 8 of 14

Phone: (215) 568-3100

8/14

503 Y S N F K Y T N A S K N P N K G V G V T TTAAATAATCTCGCTATCACTAGTTTCGTAAGGCGGAATTTAGAGGATAAACTAACCACT N N A I T SFV RRNL C D K GAATTGGTTGGAAAAACTTTAAACTTCAATAAAGCTGTAGCTGACGCTAAAAACACAGGC 583 E L V G K T L NFNK AVA D Α K CATTTAGAGAAAGAAGTAGAGAAAAAATTGGAGAGCAAAAAGCGGCAACAAAAATAAAATG 623 H L E K E V E K K L E S K S 6 GCTAATAGAGACGCAAGAGCAATCGCTTACGCTCAGAATCTTAAAGGCATCAAAAGGGAA RDARAIAYA Q N L K GAATTCAAAAATGGCAAAAATAAGGATTTCAGCAAGGCAGAAGAAACACTAAAAGCCCTT K N K D S K A E Ε AATGCAGCTTTGAATGAATTCAAAAATGGCAAAAATAAGGATTTCAGCAAGGTAACGCAA 743 N A A L N E F K N G K K V AAAGTTGATAATCTCAATCAAGCGGTATCAGTGGCTAAAGCAACGGGTGATTTCAGTAGG 783 K V D N L NGAV S ٧ Α K Α T G D CAAAAAAATGAAAGTCTCAATGCTAGAAAAAATCTGAAATATATCAATCCGTTAAGAAT 823 Q K N E S L N A RKKS Ε Ι Y Q S AAAAACTTTTCGGACATCAAGAAAGAGTTGAATGCAAAACTTGGAAATTTCAATAACAAT NFSDIKKEL AKL N N FIN G CAAGCAGCTAGCCTTGAAGAACCCATTTACGCTCAAGTTGCTAAAAAGGTAAATGCAAAA 903 Q A A S L E E ATQ V A KKVNAK CCTTTGAAAAGGCATGATAAAGTTGATGATCTCAGTAAGGTAGGGCTTTCAAGGAATCAA 943 P L K R H D D S K VI G D S TTTGGCAATCTAGAGCAAACGATAGACAAGCTCAAAGATTCTACAAAACACAATCCCATG 983 F G N EQT I D KL KDSTKHN

FIG. 4C

	_		•			_		_	_	••					_				_	
N	G	V	S	Н	L	_	٧		F			٧		I	F	N	L	Р	D	
		AL	_	_						_	IAI				Ш			-	CAAA	2280
K	G	L	S	P	Q	E	A	N	K	L	I	K	D	F	L	S	S	N	K	
		TG		AAGT							TCI				TCT		_	ACG	AGAG	2400
N	Y	D	Ε	V	K	K	A	Q	K	D	L	Ε	K	S	L	R	K	R	Ε	
GA	\GC	AAA	AG(TCA	AGC	TAA				_	_	AGAT		TGC	GTT	GAT	• • • • •		AGAG	2520
Ε	Α	K	A	Q	Α	N	S	Q	K	D	E	I	F	A	L	I	N	K	Ε	
TT	STC	TGA	\TAA	ACT	TGA	AAA	TGT	CAA	CAA	GAA	III	TGAA	IAGA	CTT	TGA	TAA	ATC	TTT	TGAT	2640
L	S	D	K	L	E	N	٧	N	K	N	L	K	D	F	D	K	S	F	D	
AAA	١GG	TTO	GGT	IGAA	AGA	TTT	AGG	TAT	CAA	TCC	AGA	AATG	GAT	TTC	AAA	AGT	TTGA	AAA	CCTT	2760
K	G	S	٧	K	D	L	G	I	N	P	E	W	I	S	K	٧	Ε	N	L	
GCA	AA	AAG	CGA	CCT	TGA	AAA	TTC	CGT	TAA	AGA	TGT	IGAT	CAT	CAA	TCA	AAA	GGT	AAC	GGAT	2880
Α	K	S	D	L	Ε	N	S	٧	K	D	٧	I	I	N	Q	K	٧	T	D	
GTA	\GA	GCA	AGC	GTT	AGC	CGA	TCT	CAA	AAA	TTT	CTO	CAAA	GGA	GCA	ATT	GGC	CCA	ACA	AGCT	3000
٧	Ε	Q	A	L	Α	D	L	K	N	F	S	Κ	Ε	Q	L	A	Q	Q	Α	
GGT	GT	GAA	TGG	AAC	CCT	AGT	CGG	TAA	TGG	GTT	AT(CTCA	AGC	AGA	AGC	CAC	CAAC	TCT	TTCT	3120
G	٧	N	G	Т	. L	٧	G	N	G	L	S	Q	Α	Ε	A	T	T	L	S	
_	AA:	TAA	TGG	ACT	CAA	AAA	_	ACC	_	TTA	_	TAA	AGT	TAA	TAA	AAA	GAA	AGC	AGGG	3240
N	N	N	G	1	K		E			Υ			٧	N			K	A	G	
			_	CAA			_			_									CTTC	3360
1	D	R	1	N	Q	1	A	S	G	, , , ,	G	۷.	ν.	G	Q	Δ	Δ	G	F	7700
GA	_	• •	TC/	• • •	_	TGA	• • •	_	_	TCA			-	_	_	TA 6	AGC	_		3480
E		A	. ι C.	K	1	D	N	1	N	Q	A	, 00. V	S	E	A	K	A	G	F	7400
_	L CT	AT/	•	•••	AAG	_	• • •	_		_		-	_	_	• • •	• • •	• • •	•	CAAT	3600
	LI								_			SIAC	1 1 1	S			1761	AGA D	M	0000
N	. ا م	W	. V	E	S	A	K		٧			_	L 	_	A	K	L [-	TAAC	7700
AAA	160	UA(CAI	UL I	AAC	.GCA	AAA	AAA	1111	. I G/	1010	ひしし	CAA	10L I	CUI	GAA	IGA	TAAG	3720

FIG. 4D

Docket No.: CHIR-0337
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Inventors: Covacci et al. Attorney: Felicity E. Groth Sheet 10 of 14

Phone: (215) 568-3100

10/14

1023 Y A T N S H I R I N S N I K N G A I ATAGTTGCGCATAATGTAGGAAGCGTTCCTTTGTCAGAGTATGATAAAATTGGCTTC 1063 I V A H N V GSVPLSEYDKI GTAAAAGACACTAATTCTGGCTTTACGCAATTTTTAACCAATGCATTTTCTACAGCA 1103 V K D T N S GFTQFLTNAF S T GGTTTCCAAAAATCTTAAAGGATTAAGGAATACCAAAAACGCAAAAACCACCCCTTG 1143 G F Q K S TGAATGCTACCAATTCATGGTATCATATCCCCATACATTCGTATCTAGCGTAGGAAG AACTCTGTAAAATCCCTATTATAGGGACACAGAGTGAGAACCAAACTCTCCCTACGG GACAGACACTAACGAAAGGCTTTGTTCTTTAAAGTCTGCATGGATATTTCCTACCCC GAAAAATCAGAAAAACCATAGGAATTATCACACCTTATAATGCCCAAAAAAGACGCT ATGCCTTTCAAGGTGAAGAGGCAGATATTATTATTATTCCACCGTGAAAACTTGTG ATCTCATTTTTGTGGGTAAAAAGTCTTTCTTTGAGAATTTATGAAGCGATGAGAAGA CATTCTTCGCTTCAAAACGCTTTCATAAATCTCTCTAAAGCGCTTTATAATCAACAC TTATTAGCGTTACAATTTGAGCCATTCTTTAGCTTGTTTTTCTAGCCAGATCACATC TAAAATAATCACTTCGGGAAAATCTTTAAGGGAGTGAAATAATAACGCATGCAAGTT TGCGAAACATTCAAATAGCCTTGTTGTTTCAGGGCATTGTCATAAGCGTTGGATTGG GCTAAAATGCTTGGCTCAATCACGCCCACAATAGGGATTTTGGAATGCTTTTGCATC TTGAAAAATCCAAAGCCTCTAAGCCAAATTGCTTGATCGTAGTGGGGTCTTTAGTG **AGGCTTTTTAAAACGCTAAACCCTCCCACACCGCTATCAAAAACGCCTATTTTCATG** TCTTCATTGTCCTTAGTTTGTTGCATTTTAGAATAGACAAAGCTT 5925

FIG. 4E

Ε	K	A	١	Т	G	M	L	Т	Q	K	٨	ł	Р	E	W	1	K		v	M			
AA	CCA	AGA	AG	ΔΔ	ΓΔΤ	SAS	AGA	TT	TT					~~				ا امد د	V	N	D	K	
N	Q	K	,	N	M	u v	707			C 1 G	M I I	ָנט					.CA(CAA	IGTT	GAA	CAA	TGCT	3840
	_	• • •	•	••	• •	K	D	Υ	S	D	S		F	K	F	S	T	K	L	N	N	Α	
10	117	111	AC	160	:110	GC	GAG	AGA	AA	ATG	CGG	AG	CA.	TGG	AAT	CAA	GAA	CGT	TAA	TAC	AAA	AGGT	3960
2	Y	Y		C	L	Α	R	Ε	N	Α	Ε		Н	G	I	K	N	V	N	Т	V	G	7300
CI	AAA	AG	CG	AGG	GGT	TT.	TTT	AAT	ACT	CCT	ΓTΑ	GC	AG	AAA	TCC	CAA	TCG	TCT	TTA	GTA	ΙΊΤ	GGGA	4080
TG	TGC	AA	AG	TTA	CGC	CT	TTG	GAG	ΔΤΔ	TG	ATG	TG	TG	SCA	CCT	GTA	ccc	AAT		T TA		CTCA	
GC	ΔΔς	ΔΤ	CA	crr	TAG	GA	c	CCA	ATC	·	110 TT	TA	1 U.	107		GIA	900	AAI	666	116	GAG	CTCA	4200
AA	AAA	CA	CT		CCC	TAD	100	ししみ	AIL	.610	-	IA	5 C	961	IGG	GCA	CTT	CAC	CTT	AAA	ATA'	TCCC	4320
AA	***	GAI		IAA	ししし	. 1 1 1	GC	ITA	AAA	TTA	IAG	П	TGA	TT(GTG	CTA	GTG	GGT	TCG	TGC	TAT	AGTG	4440
AL	AAA	GA	I C	AAG	TTC	AAA	IAA/	ATC	ATA	GAG	ict'	TT	TAG	AG	CAA	ATT	GAT	CGC	GCT	CTT	445	CAAA	4560
TG	CGA	TC	AG/	AAG	TGG	AAA	AA	TAC	GGC	TTC	AA	GA/	ATT	TT	SAT	SAG	CTC	ΔΔΔ	ΔΤΑ	GAC	ACT	GTGG	
GT	TAP	CT	TT(TT	TCT	TGC	TAG	AT	TCT	ΔΔΔ	re	CTT	ΓGA	ATO	TC	CT	ATT	TCT	A C C	CCA	40 I (SAAA	4680
AT	TC:	TT	ΓΔί	ire	CTA	TTT	TG	• A A	CTC	TOT	AC	AT		744	3 I G(36 I	M	161	AUU	SCA/	AAA	AAA	4800
ΔΔ1		ררז		TA	CTC	1 1 1 TCA	201	, A A .	916	101	4U/	41/	100	IAA	AIC!	111	I CC.	AAA	GAT	AAT	CATI	TAGA	4920
CC				IA	916	IGA	661	AI	460	CCC	П	П	GG	GAA	TT	SAG	TTA'	ITT.	TGA	CTT	ΓΑΑ	ATTT	5040
GC	.66	C	CA	TG	AAA	TTC	CAC	TT:	TAG	GGA	AT(GC (iTG	TGC	CATT	TT	ITT	ΓΑΑ(3GG(CGTA	TTI	TTTG	5160
GG(:AA	TAP	GC	:TC	CAT	AAA	ATA	GC	CT	CAA	TTI	Ш	TG	AGC	GAT	TA	AGG	SAA	ATO	CGT	GCA	ACC	5280
TCT	'AA(CAA	TT	CG	CCC	TCT	AAA	AT	CT	TTC	TTC	`ΔΔ	TC	ΔΔΔ	GGC	ACI	MAA	AG	SGA	CTO	CCT	TAAA	. –
ATC	ĠT(GC	TT	TT	iTC	CCT	AGC	AC1	ΓΔ Δ.	AAT	AGG	:66		TTT	TT4	TC1	100/	140/	TC.	1010	100	SATC	5400
TCT	Tri	ΓΔΑ	AG	CT	S C A		MUU CTC		CAT!	RAI.	70°	700		111	IIA	116		AU	161	CGC	TTG	ATC	5520
	TA		70	CT	AUA	JUU	いし	ו טט.	61	011	6C/	١IG	CC	ACA	ATC	AA	TAA7	TCA	NAT(TGG	TGC	GGT	5640
LLA	IAA	106	LA	CI(. I A	CC	GTA	TC	iCC/	ATA	ATA	IGA	TG	ATT	TCA	TCA	IAAI	'AA1	TGC	GCT	TTT	AAA	5760
ACA	CTT	П	П	AAT	TT	TAP	GGG	ATT	'AA'	TTA	GGG	AT	TT	TAT	TTT	TCA	TTC	AT1	AAG	ITT	AAA	AAT	5880

FIG. 4F

Docket No.: CHIR-0337

App No.: Not Yet Assigned Filed: Herewith Title: HELICOBACTER PYLORI CAI ANTIGEN
Inventors: Covacci et al.
Attorney: Felicity E. Groth Phone: (215) 568
Sheet 12 of 14

Phone: (215) 568-3100

	10 30 50 AAGCTTGCTGTCATGATCACAAAAAACACTAAAAAACATTATTATT <u>AAGGA</u> TACAAAATG																			
A	AGC	TT	GC1	GT	CA.	TGA	TCA	CAA	AAA.	ACA	CTA	AAA	AAC	ATT	ATT.	ATT	AAG	GAT	ACA	AAATG
																		•		М
				0							90						110	0		
G	CAA	AA	GAA	AT	CA	TAP	III	CAG	ATA	GTG	CGA	GAA	ACCI	П	TAT	TTG	AAG	scg	TGA	GGCAA
A	K		E	I	K				S					L	_	Ε		٧	R	
			13	-							50						170) ·	•••	
C.	TCC	AT	SAC	GC	TGT	CA	AAG"	TAA	CCAT	TGG	GGC	CAAG	SAGO	CAG	GA/	ATG	TAT	ΓGA	TCC.	AAAA
L	Н			A	٧	K			M		P			R		٧	L	I	Q	K
			19	-							10						230)	_	
A(SCT	ATG	GC	GC	TCC	AAG	CAT	CAC	CAA	\AG/	ACG(GCGT	GAG	CGT	GGC	TA	AAGA	IGA"	TTG	AATTA
S	Y			A	P	S	I		K		G		S				Ε	I	Ε	L
			25	-						27							290)		_
AG	iTT(GCC	CA	GTA	IGC	TAA	CAT	GGG	CGC	TCA	ACT	CGT	TAA	AGA	AGT	AG	CGAG	CAA	AAA(CCGCT
S	C	P	, 1	/	A	N	M		A		L		K		٧	A	S	K	T	A
			310	-						33							350)		
GA	TG(CTG	CCG	GC	GA	TGG	CAC	GAC	CAC	AGC	GAC	CGT	GCT	AGC	TTA	TAG	CAT	TTI	TAA	AGAA
D	A	A			D	G			T			٧				S	I	F		Ε
			370							39	_						410			
GG	TTI	TGA	GGA	AT	AT	CAC	GGC	TGG	GGC	TAA	CCC	TAT	TGA.	AGT	GAA	ACG	AGG	CAT	GGA	TAAA
G	L	R	N	1	I	T	A	G	A	N	P	I	Ε	٧	\mathbf{K}^{\cdot}	R	G	M	D	K
			430							45							470			
GC	TGC	TG	AAG	CG	AT(CAT	TAA	TGA	GCT	TAA	AAA	AGC	GAG	CAA	AAA	AGT	AGG	CGG	TAA	AGAA
A	A	E	A	1	I	I	N	Ε	L	K	K	A	S	K	K	٧	G	G	K	Ε
			490							51							530			
GA	AAT	CA	CCÇ	AA	GTO	GC	GAC	CAT	TTC	TGC	AAA	CTC	CGAT	ГСА	AA'	TAT	CGG	GAA	ACT	CATC
E	I	T	0		V	A	T	I	S	A		S	D	H	N	I	G	K	L	I
		_	550							570							590		-	_
GC.	TGA	CG	CTA	TG	GAA	AA	AGTO	GGG	ΓΑΑ	AGA	CGG	CGT	SATO	CACC	GTT	ΓGA	GGA	AGC	TAA	GGGC
A	D	A	M		E	K	٧	G	K			٧		T		Ε			K	
			510							630	-						650			-
	ΓGA.	AGA	\TG	AAT	TTG	GAT	TGT(CGTA	\GA/	AGG(CAT	GCAA	ITT	GAT	AGA	AGG	CTAC	CT	CTC	CCCT
I	Ε	D		į		D	٧		Ε	G	M		F	D	R	G	Y		S	Р

FIG. 5A

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13/14 670 690 710 TATTTTGTAACGAACGCTGAGAAAATGACCGCTCAATTGGATAATGCTTACATCCTTTTA YFVTNAEKMTAQLDNAYILL *7*30 750 770 ACGGATAAAAAATCTCTAGCATGAAAGACATTCTCCCGCTACTAGAAAAAACCATGAAA T D K K I S S M K D I L P L L E K T M K 790 810 HindIII GAGGGCAAACCGCTTTTAATCATCGCTGAAGACATTGAGGGCG<u>AAGCTT</u>TAACGACTCTA GKPLL EDIEGEALTTL Ī I A 850 870 890 GTGGTGAATAAATTAAGAGGCGTGTTGAATATCGCAGCGGTTAAAGCTCCAGGCTTTGGG V V N K L R G V L N I A A V K A P G F G 910 930 950 GACAGAAGAAAGAAATGCTCAAAGACATCGCTATTTTAACCGGCGGTCAAGTCATTAGC DRRKEMLKDIAILTGGQVIS 970 990 1010 GAAGAATTGGGCTTGAGTCTAGAAAACGCTGAAGTGGAGTTTTTAGGCAAAGCTGGAAGG E E L G L S L'EN A E V E F L G K A G R 1030 1050 1070 ATTGTGATTGACAAAGACAACACCACGATCGTAGATGGCAAAGGCCATAGCGATGATGTT IDKDNTTIVDGKGHSDDV 1090 1110 1130 AAAGACAGAGTCGCGCAGATCAAAACCCAAATTGCAAGTACGACAAGCGATTATGACAAA RVAQIKTQIASTTSDYDK 1150 1170 1190 GAAAAATTGCAAGAAAGATTGGCTAAACTCTCTGGCGGTGTGGCTGTGATTAAAGTGGGC EKLQERLAKLS G G V A V I K V G 1210 1230 1250 GCTGCGAGTGAAGTGGAAATGAAAGAGAAAAAAGACCGGGTGGATGACGCGTTGAGCGCG A A S E V E M K E K K D R V D D A L S A 1270 1290 1310 ACTAAAGCGGCGGTTGAAGAAGGCATTGTGATTGGTGGCGGTGCGGCTCTCATTCGCGCG TKAAVEEGIVIGGGAALI

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14/14

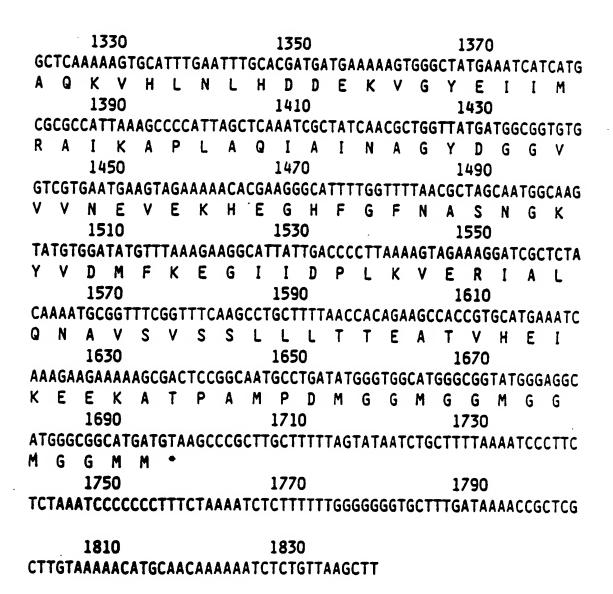


FIG. 5C